Entry Page Page 1 of 1



## Text Entry | SwissEntry

```
Reset
                                                   General Description References Links Sequence
       Entry
   Information
                         General information
                                          IPI00083978
                         Entry name
· Entry from:
V
                         Accession number IPI00083978.2
  Entry Options
                        Created
                                          IPI HUMAN Rel. 2.5, 4-MAR-2003
                         Sequence update IPI HUMAN Rel. 2.8, 6-JUN-2002
Launch analysis tool:
                         Description and origin of the Protein
BlastP
                         Description
                                          SIMILAR TO CD63 PROTEIN.
            Launch
                         Organism source
                                          Homo sapiens (Human).
                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Link to related
                        Taxonomy
information:
                                          Primates; Catarrhini; Hominidae; Homo.
              Link
                         NCBI TaxID
                                          9606
                         Database cross-references
Save entry:
              Save
                         InterPro
                                          <u>IPR000301</u>; Transmem_4.
View:
                         Pfam
                                          PF00335; transmembrane4; 1.
 Printer Friendly
                         PROSITE
                                          PS50257; TM4_2; 1.
                         REFSEQ_XP
                                          XP_084868; GI:20553887; M.
                         Sequence information
                         Length: 220 aa, molecular weight: 25225 Da, CRC64 checksum: 551BC6E4A
                              MGFGAWLLLD RNNFLTAFDE NNHFIVPISQ ILIGMGSSTV LFCLLGYIGI HNEIRWL:
                              YAVLITWTFA VQVVLSAFII TKKEEVQQLW HDKIDFVISE YGSKDKPEDI TKWTILN
                              KTLQCCGQHN YTDWIKNKNK ENSGQVPCSC TKSTLRKWFC DEPLNATYLE GCENKIS.
                              NVNVLTLIGI NFGLLTSEVF QVSLTVCFFK NIKNIIHAEM
                         11
```

General Description References Links Sequence

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